

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/537,017  
Source: PCT  
Date Processed by STIC: 6-13-05

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PCT

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/10/537,017**

**DATE: 06/13/2005**  
**TIME: 09:40:56**

**Input Set : A:\67314-02 SEQ LIST.txt**  
**Output Set: N:\CRF4\06132005\J537017.raw**

3 <110> APPLICANT: THE STATE OF OREGON ACTING BY AND THROUGH THE STATE BOARD  
 4 OF HIGHER EDUCATION ON BEHALF OF OREGON STATE UNIVERSITY  
 5 Azevedo, Mark  
 6 Armstrong, Donald  
 7 Mills, Dallice I.  
 8 Banowetz, Gary  
 9 Russell, Brian  
 10 Groenig, Aleta  
 11 Elliott, Lloyd  
 13 <120> TITLE OF INVENTION: BACTERIAL BIOHERBICIDE FOR CONTROL OF GRASSY WEEDS  
 15 <130> FILE REFERENCE: 245-67314-02  
 C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/537,017  
 C--> 17 <141> CURRENT FILING DATE: 2005-06-01  
 17 <150> PRIOR APPLICATION NUMBER: US 60/431,651  
 18 <151> PRIOR FILING DATE: 2002-12-06  
 20 <150> PRIOR APPLICATION NUMBER: PCT/US2003/038653  
 21 <151> PRIOR FILING DATE: 2003-12-05  
 23 <160> NUMBER OF SEQ ID NOS: 13  
 25 <170> SOFTWARE: PatentIn version 3.2  
 27 <210> SEQ ID NO: 1  
 28 <211> LENGTH: 550  
 29 <212> TYPE: DNA  
 30 <213> ORGANISM: Pseudomonas fluorescens  
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 37 ggccagtgtc aactgtcgat cctgttgcgt gtattggccgg gcgagcgatc cgtggcgccg 180  
 39 ggggtctggc gccccgggtgc ggtgttgcgtt aagttattgg tgggcagcaaa ggccggcgccg 240  
 41 cattttcagc gtgaactcac gggcgtgcgc ctgtggccgg aacacggcct gaccaccccc 300  
 43 cgggtgtcgcc ccgtatggctt gcaggaaaggc gagggcggtt gttgtctgtt cgagtttcctc 360  
 45 gaaggccggc aaaggcttgc cgtatgcctgg caggcccggtt cgaagcgctg ccgcggctgg 420  
 47 ccgacgaaca aaccgcgggtg ctgcggcaag cgctgggtgc gatcgccgac atgcacacca 480  
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 51 atcgacgtgc 550  
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 64 aagcacttggc ggcgtggggcc aggtgggtgc caaccgggttc gatcttgatt ttcttcggat 180  
 66 cgaaggccag ttgctccagg caccagacga tcttggcgat gtagcgcttg tagcgacgg 240

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72	cggaaaggctt	gatgcggcc	tgttccaggc	agaacttcgc	cgactcgtag	420	
74	tctttgcatg	tttgcgtcg	acgaagcgt	cttcttcggc	ggccgcgatc	480	
76	cgcataatacg	ggctgcggaa	ggatcatggc	taagggcgc	ggacaggcca	540	
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84	cactttcccg	gtagatcgta	taggggagaa	tgagcatgtc	gaggacgcgg	780	
86	agtcaagaac	ggcccagggc	acgcggggaa	tgaatccgt	aatacgtatcg	840	
88	cgcgtggaa	cgcataatac	aggcctctca	cgtgaagcac	ggggaaaccat	cacacttgc	900
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92	accgcgcggc	atccctggca	tatgcgggt	cgaaactggc	ggctgagctg	tgctgacaga	1020
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116	cgagcaggt	atacacc	agctcctcg	taaacgg	gaggttt	gcaatttgg	1740
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120	cgacacggat	gcccgcacc	tcgatcaat	acagcttgc	gtcctggc	agcagg	1860
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124	ttcggc	gag	ccaccgg	tgttgc	ccagcgg	cagcg	1980
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130	gcacccc	gtat	ccac	tgaaat	gcg	cccacc	2160
132	ccagcacc	acg	cccg	cagacc	cc	gcaata	2220
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144	gggtt	cagg	tcg	act	cg	acc	2580
146	tatgtt	gg	taaaa	acag	acc	tgca	2640
148	gccac	agg	cagc	aggat	ggg	gtcg	2700
150	gcacc	aa	cagg	tc	cc	caac	2760
152	cgcc	agg	cc	tcg	cc	gcgt	2820
154	gata	acgg	at	tgc	cc	ccgt	2880
156	gact	cg	aa	cgc	gg	ccgt	2940
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247				20			25				30				
250 Leu	Ser	Leu	Leu	Arg	Val	Leu	Pro	Gly	Glu	Arg	Tyr	Val	Gly	Ala	Ala
251				35			40				45				
254 Val	Trp	Arg	Gly	Arg	Ala	Val	Leu	Ala	Lys	Leu	Leu	Val	Gly	Ser	Lys
255				50			55				60				
258 Ala	Ala	Arg	His	Phe	Gln	Arg	Glu	Leu	Thr	Gly	Val	Arg	Leu	Leu	Ala
259				65			70				75			80	

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263 85 90 95  
266 Gly Glu Gly Gly Trp Leu Leu Phe Glu Phe Leu Glu Gly Ala Glu Ser  
267 100 105 110  
270 Leu Ala Asp Ala Trp Gln Ala Val Glu Ala Leu Pro Pro Leu Ala Asp  
271 115 120 125  
274 Glu Gln Thr Ala Val Leu Ala Glu Ala Leu Gly Ala Ile Ala Gln Met  
275 130 135 140  
278 His Thr Lys Gly Leu Trp Gln Glu Asp Leu His Leu Asp Asn Leu Leu  
279 145 150 155 160  
282 Arg Gln Asp Gly Lys Leu Tyr Leu Ile Asp Gly Ala Gly Ile Arg Val  
283 165 170 175  
286 Glu Glu Ala Gly Lys Pro Leu Ser Arg Asn Arg Val Leu Glu Asn Leu  
287 180 185 190  
290 Gly Val Phe Phe Ala Gln Leu Pro Lys Asn Leu Glu Pro Phe Thr Glu  
291 195 200 205  
294 Glu Leu Leu Val Tyr Tyr Leu Leu Gly Asn Ser Glu His Ala Leu Pro  
295 210 215 220  
298 Leu Gln Ala Leu Glu Lys Gln Val Arg Lys Val Ser Ala Trp Arg Leu  
299 225 230 235 240  
302 Lys Asp Phe Leu Asn Lys Ala Gly Arg Glu Cys Thr Leu Phe Ser Val  
303 245 250 255  
306 Val Arg Gly Ala Phe Gly Leu Arg Ala Ile Arg Arg Glu Glu Glu Pro  
307 260 265 270  
310 Ala Met Leu Pro Val Leu Glu Gln Ala Asp Ala Leu Leu Asp Arg Gly  
311 275 280 285  
314 His Leu Tyr Lys Thr Gly Gly Ala Ala Ser Val Ala Lys Val Glu Val  
315 290 295 300  
318 Ala Gly Arg Pro Leu Val Ile Lys Arg Tyr Asn Ile Lys Gly Phe Ala  
319 305 310 315 320  
322 His Trp Leu Lys Arg Phe Trp Arg Pro Ser Arg Ala Trp His Ser Trp  
323 325 330 335  
326 Arg Glu Gly Asn Arg Leu Ala Phe Leu Gly Ile Ala Thr Pro Lys Pro  
327 340 345 350  
330 Leu Ala Val Leu Glu Lys Arg Val Phe Trp Leu Arg Ser Arg Ala Tyr  
331 355 360 365  
334 Leu Ile Thr Glu Tyr Leu Pro Gly Pro Asp Ile Ile Glu Arg Phe Ala  
335 370 375 380  
338 Pro Tyr Ile Asp Asn Gly Asp Ala Pro Glu Asn Glu Leu Leu Ala Leu  
339 385 390 395 400  
342 Asp His Leu Phe Ala Gln Leu Ile Gly Glu Arg Ile Ser His Gly Asp  
343 405 410 415  
346 Phe Lys Gly His Asn Leu Phe Trp His Glu Asp Arg Trp Ala Leu Ile  
347 420 425 430  
350 Asp Leu Asp Ser Val Cys Gln His Ser Ser Ala Ala Ser Phe Ala Pro  
351 435 440 445  
354 Ala Tyr Ala Arg Asp Arg Ala Arg Phe Gln Val  
355 450 455  
358 <210> SEQ ID NO: 4

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 361 <213> ORGANISM: Pseudomonas fluorescens  
 363 <400> SEQUENCE: 4  
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 374 35 40 45  
 377 Thr Ala Ala Gln Pro Glu Asn Ala Leu Phe Gln Tyr Gly Gln Trp Leu  
 378 50 55 60  
 381 Gly Arg Gly Asp Thr Gln Glu Arg Gln Ala Val Ala Phe Thr Pro Gly  
 382 65 70 75 80  
 385 Val Pro Gly Ala Ala Trp Ala Pro Glu Ala Phe Glu Pro Met Gly Glu  
 386 85 90 95  
 389 Pro Asp Val Val Thr Leu Asp His Gln Arg Thr Thr Gly His Leu Asp  
 390 100 105 110  
 393 Leu Gly His Ala Gly Gly Thr Ala Gly Leu Val Gln Val Ala Ala Val  
 394 115 120 125  
 397 Glu Gln Arg Ile Gly Leu Phe Gln His Arg Gln His Arg Arg Leu Phe  
 398 130 135 140  
 401 Phe Ala Thr Asn Arg Ala Gln Thr Lys Arg Thr Ala His Tyr Ala Glu  
 402 145 150 155 160  
 405 Gln Arg Ala Phe Thr Ala Gly Leu Val Gln Lys Val Leu Gln Ala Pro  
 406 165 170 175  
 409 Gly Asp Leu Ala His Leu Phe Phe Gln Arg Leu Gln Arg Gln Arg Val  
 410 180 185 190  
 413 Leu Thr Val Ala Glu Gln Val Ile His Gln Gln Leu Leu Gly Lys Arg  
 414 195 200 205  
 417 Leu Glu Val Phe Arg Gln Leu Gly Lys His Thr Glu Val Phe Gln Tyr  
 418 210 215 220  
 421 Ser Val Ala Arg Gln Arg Phe Thr Gly Leu Phe Thr Asp Ala Arg Thr  
 422 225 230 235 240  
 425 Val Asp Gln Val Gln Leu Ala Val Leu Ala Gln Gln Val Val Gln Met  
 426 245 250 255  
 429 Gln Val Phe Leu Pro Gln Pro Phe Gly Val His Leu Arg Asp Arg Thr  
 430 260 265 270  
 433 Gln Arg Phe Gly Glu His Arg Gly Leu Phe Val Gly Gln Arg Arg Gln  
 434 275 280 285  
 437 Arg Phe Asp Gly Leu Pro Gly Ile Gly Gln Ala Phe Gly Ala Phe Glu  
 438 290 295 300  
 441 Glu Leu Glu Gln Gln Pro Thr Ala Leu Ala Phe Leu Gln Ala Ile Gly  
 442 305 310 315 320  
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RAW SEQUENCE LISTING ERROR SUMMARY                    DATE: 06/13/2005  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 444

Seq#:7; N Pos. 691

**VERIFICATION SUMMARY**

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Input Set : A:\67314-02 SEQ LIST.txt  
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L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:420

L:605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:660